

SEQUENCE LISTING

5 <110> Herr, John C.
Norton, Elizabeth J.
Deikman, Alan B.

10 <120> Recombinant Antibody Directed Against Human Sperm
Antigen

<130> 00415-02

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<150> 60/145,512
<151> 1999-07-23

20 <160> 18

<170> PatentIn Ver. 2.1

<210> 1
25 <211> 116
<212> PRT
<213> Mus musculus

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Gly Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser
20 25 30

35 Asn Arg Asp Thr Tyr Leu His Trp Phe Leu Gln Lys Pro Gly Gln Ser
35 40 45

Pro Glu Leu Leu Ile Tyr Arg Val Ser Asn Arg Phe Ser Gly Val Pro
40 50 55 60

Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
65 70 75 80

Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Phe Cys Ser Gln Ser
85 90 95

5 Thr His Val Pro Phe Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys
100 105 110

Arg Ala Ala Ala
115

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<210> 2
<211> 348
<212> DNA
15 <213> Mus musculus

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20 atctcttgca gatctagtca gagtcttgta cacagtaata gagacactta ttacattgg 120
ttcctgcaga agccaggcca gtctccagag ctctgatct acagagtttc caaccgattt 180
tctgggggtcc cagacagggt cagtggcagt ggatcagga cagatttcac actcaagatc 240
25 agcagagtgg aggctgagga tctgggagtt tatttctgtt ctcaaagtac acatgttcca 300
ttcacgttcg gctcggggac caagctggaa atcaaacggg cggccgca 348

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<210> 3
<211> 118
<212> PRT
35 <213> Mus musculus

<400> 3
Gln Val Lys Leu Gln Gln Pro Gly Ser Glu Pro Val Arg Pro Gly Ala
1 5 10 15
40 Ser Val Lys Val Ser Cys Arg Ala Ser Gly Tyr Lys Phe Thr Thr Tyr
20 25 30
Trp Met His Trp Val Arg Gln Arg Pro Gly Gln Gly Pro Glu Trp Ile
45 35 40 45

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Gly Asp Ile Tyr Pro Gly Ser Gly Asp Ser Asn Tyr Asp Val Lys Phe
 50 55 60

5 Lys Asn Lys Ala Thr Leu Thr Val Asp Thr Ser Ser Ser Thr Val Tyr
 65 70 75 80

Ile Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
 85 90 95

10 Ala Arg Gly Asp Tyr Gly Cys Pro Phe Val Tyr Trp Gly Gln Gly Thr
 100 105 110

Thr Val Thr Val Ser Ser
 15 115

<210> 4
 <211> 354
 20 <212> DNA
 <213> Mus musculus

<400> 4
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 25 tcctgcaggg cttctggcta caaattcacc acctactgga tgcactgggt gaggcagagg 120
 cctggacaag gccctgagtg gattggagat atttatcctg gtagtgggtga ttctaactac 180
 30 gatgtgaagt tcaagaacaa ggccacactg actgtagaca catcctccag cacagtttac 240
 atacaactca gcagcctgac atctgaggac tccgcggtct attactgtgc aagagggggac 300
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<210> 5
 <211> 15
 40 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:peptide linker
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<400> 5
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<210> 6
 <211> 100
 <212> DNA
 10 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:PCR primer

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 ggcggcagcg acatcgagct cactcagtct ccattctccc 100
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<210> 7
 <211> 100
 <212> DNA
 25 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:PCR primer

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 cgctgccgcc gccgccactg gagacggtga ccgtggtgcc 100
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<210> 8
 <211> 264
 <212> PRT
 40 <213> Mus musculus

<400> 8
 Met Ala Gln Val Lys Leu Gln Gln Pro Gly Ser Glu Pro Val Arg Pro
 1 5 10 15
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Gly Ala Ser Val Lys Val Ser Cys Arg Ala Ser Gly Tyr Lys Phe Thr
 20 25 30

5 Thr Tyr Trp Met His Trp Val Arg Gln Arg Pro Gly Gln Gly Pro Glu
 35 40 45

Trp Ile Gly Asp Ile Tyr Pro Gly Ser Gly Asp Ser Asn Tyr Asp Val
 50 55 60

10 Lys Phe Lys Asn Lys Ala Thr Leu Thr Val Asp Thr Ser Ser Ser Thr
 65 70 75 80

Val Tyr Ile Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
 15 85 90 95

Tyr Cys Ala Arg Gly Asp Tyr Gly Cys Pro Phe Val Tyr Trp Gly Gln
 100 105 110

20 Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly
 115 120 125

Gly Ser Gly Gly Gly Gly Gly Ser Asp Ile Glu Leu Thr Gln Ser Pro Phe
 130 135 140

25 Ser Leu Pro Val Ser Leu Gly Gly Pro Ala Ser Ile Ser Cys Arg Ser
 145 150 155 160

Ser Gln Ser Leu Val His Ser Asn Arg Asp Thr Tyr Leu His Trp Phe
 30 165 170 175

Leu Gln Lys Pro Gly Gln Ser Pro Glu Leu Leu Ile Tyr Arg Val Ser
 180 185 190

35 Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly
 195 200 205

Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly
 210 215 220

40 Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Phe Thr Phe Gly Ser
 225 230 235 240

Gly Thr Lys Leu Glu Ile Lys Arg Ala Ala Ala Gly Ala Pro Val Pro
 45 245 250 255

Tyr Pro Asp Pro Leu Glu Pro Arg
260

5

<210> 9

<211> 792

<212> DNA

<213> Mus musculus

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aaggtgtcct gcagggcttc tggctacaaa ttcaccacct actggatgca ctgggtgagg 120

15

cagaggcctg gacaaggccc tgagtggatt ggagatattt atcctggtag tgggtgattct 180

aactacgatg tgaagttcaa gaacaaggcc aactgactg tagacacatc ctccagcaca 240

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ggggactatg gttgcccttt tgtttactgg ggccaaggca ccacggtcac cgtctccagt 360

ggcggcggcg gcagcgggtgg tgggtggttct gggggcggcg gcagcgacat cgagctcact 420

25

cagtctccat tctccctgcc tgtcagtctt ggaggtccag cctccatctc ttgcagatct 480

agtcagagtc ttgtacacag taatagagac acttatttac attgggttcct gcagaagcca 540

30

ggccagtctc cagagctcct gatctacaga gtttccaacc gattttcttg ggtcccagac 600

aggttcagtg gcagtggatc agggacagat ttcacactca agatcagcag agtggaggct 660

gaggatctgg gagtttattt ctgtttctcaa agtacacatg ttccattcac gttcggctcg 720

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gggaccaagc tggaaatcaa acgggcggcc gcaggtgcgc cggtgccgta tccggatccg 780

ctggaaccgc gt 792

40

<210> 10

<211> 792

<212> DNA

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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic
sequence substituting bacterial codons for mouse
codons

5

<400> 10

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10 aaggtgtcct gccgcgcttc tggctacaaa ttcaccacct actggatgca ctgggtgcg 120
cagcgccctg gccaaaggccc tgagtggatt ggcgatatatt atcctggtag tggtgattct 180
aactacgatg tgaagttcaa gaacaaggcc aactgactg tagacacatc ctccagcaca 240
15 gtttacatcc aactcagcag cctgacatct gaggactccg cgggtctatta ctgtgcaaga 300
ggggactatg gttgcccttt tgtttactgg ggccaaggca ccacggtcac cgtctccagt 360
20 ggcgggcgcg gcagcggtgg tgggtggttct gggggcgggc gcagcgacat cgagctcact 420
cagtctccat tctccctgcc tgtcagtctt ggcgatccag cctccatctc ttgccgctct 480
agtcagagtc ttgtacacag taatcgcgac acctatctgc attgggttct gcagaagcca 540
25 ggccagtctc cagagctcct gatctaccgc gtttccaacc gcttttctgg ggtcccagac 600
cgcttcagtg gcagtggctc agggacagat ttcacactca agatcagcag cgtggaggct 660
30 gaggatctgg gcgtttatct ctgttctcaa agtacacatg ttccattcac gttcggctcg 720
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ctggaaccgc gt 792

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<210> 11

<211> 251

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<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic
sequence substituting amino acids in the natural

45

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mouse protein to "humanize" the protein

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<400> 11
Met Ala Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Val Lys Lys Pro
5      1              5              10              15

Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr
      20              25              30

10 Thr Tyr Trp Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu
      35              40              45

Trp Ile Gly Asp Ile Tyr Pro Gly Ser Gly Asp Ser Asn Tyr Asp Val
      50              55              60

15 Lys Phe Lys Asn Arg Val Thr Ile Thr Ala Asp Thr Ser Thr Ser Thr
      65              70              75              80

Ala Tyr Met Gln Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr
20      85              90              95

Tyr Cys Ala Arg Gly Asp Tyr Gly Cys Pro Phe Val Tyr Trp Gly Gln
      100             105             110

25 Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly
      115             120             125

Gly Ser Gly Gly Gly Gly Ser Asp Ile Val Met Thr Gln Ser Pro Ser
      130             135             140

30 Ser Leu Pro Val Ser Val Gly Asp Pro Ala Ser Ile Ser Cys Arg Ser
      145             150             155             160

Ser Gln Ser Leu Val His Ser Asn Arg Asp Thr Tyr Leu His Trp Tyr
35      165             170             175

Leu Gln Lys Pro Gly Gln Ser Pro Gln Leu Leu Ile Tyr Arg Val Ser
      180             185             190

40 Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly
      195             200             205

Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly
      210             215             220

45

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Val Tyr Tyr Cys Ser Gln Ser Thr His Val Pro Phe Thr Phe Gly Gln
 225 230 235 240

5 Gly Thr Lys Val Glu Ile Lys Arg Ala Ala Ala
 245 250

<210> 12
 10 <211> 753
 <212> DNA
 <213> Artificial Sequence

<220>
 15 <223> Description of Artificial Sequence: synthetic
 sequence substituting human codons for mouse
 codons

<400> 12
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 caagctcctg gtcaaggctt ggaatggatt ggtgatattt atcctgggtc tggtgattct 180
 25 aattatgatg ttaaatttaa aaatcgtgtt accattaccg ctgatactc tacctctacc 240
 gcttatatgc aattatctag cttacgttct gaagataccg cagtttatta ttgtgcacgt 300
 30 ggtgattatg gttgtccttt tgtttattgg ggtcaaggca ccacgggtac cgtttctagc 360
 ggtggcggcg gttctggcgg tggcggtagc ggcggtggtg gctctgatat tgttatgacc 420
 caatctcctt ctagcttacc tgtttctggt ggtgatcctg ctagcatttc ttgtcgttct 480
 35 agccaatctt tagttcatag caatcgtgat acctatttac attggtatct gcagaaacct 540
 ggtcaaagcc ctcaattact gatttatcgt gttagcaatc gtttttagcgg tgttcctgat 600
 40 cgtttttctg gtagcgggtc tggtagcgat ttacggttaa aaatttctcg tgttgaagct 660
 gaggatggtg gtgtttatta ttgttctcaa agcaccatg ttccttttac gttcgggtcaa 720
 ggtaccaaag ttgaaattaa acgtgctgca gct 753
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<210> 13
 <211> 45
 5 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:nucleic acid
 10 linker

<400> 13
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 15

<210> 14
 <211> 13
 <212> PRT
 20 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:commercially
 available petide antigen
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<400> 14
 Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu Glu Pro Arg
 1 5 10

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 <210> 15
 <211> 251
 <212> PRT
 <213> Mus musculus

35
 <400> 15
 Met Ala Gln Val Lys Leu Gln Gln Pro Gly Ser Glu Pro Val Arg Pro
 1 5 10 15

40 Gly Ala Ser Val Lys Val Ser Cys Arg Ala Ser Gly Tyr Lys Phe Thr
 20 25 30

Thr Tyr Trp Met His Trp Val Arg Gln Arg Pro Gly Gln Gly Pro Glu
 35 40 45
 45

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Trp Ile Gly Asp Ile Tyr Pro Gly Ser Gly Asp Ser Asn Tyr Asp Val
 50 55 60

5 Lys Phe Lys Asn Lys Ala Thr Leu Thr Val Asp Thr Ser Ser Ser Thr
 65 70 75 80

Val Tyr Ile Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
 85 90 95

10 Tyr Cys Ala Arg Gly Asp Tyr Gly Cys Pro Phe Val Tyr Trp Gly Gln
 100 105 110

Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly
 15 115 120 125

Gly Ser Gly Gly Gly Gly Ser Asp Ile Glu Leu Thr Gln Ser Pro Phe
 130 135 140

20 Ser Leu Pro Val Ser Leu Gly Gly Pro Ala Ser Ile Ser Cys Arg Ser
 145 150 155 160

Ser Gln Ser Leu Val His Ser Asn Arg Asp Thr Tyr Leu His Trp Phe
 165 170 175

25 Leu Gln Lys Pro Gly Gln Ser Pro Glu Leu Leu Ile Tyr Arg Val Ser
 180 185 190

Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly
 30 195 200 205

Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly
 210 215 220

35 Val Tyr Phe Cys Ser Gln Ser Thr His Val Pro Phe Thr Phe Gly Ser
 225 230 235 240

Gly Thr Lys Leu Glu Ile Lys Arg Ala Ala Ala
 245 250

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<210> 16
 <211> 753
 <212> DNA
 45 <213> Mus musculus

-12-

<400> 16
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 cagaggcctg gacaaggccc tgagtggatt ggagatattt atcctggtag tggtgattct 180
 aactacgatg tgaagttcaa gaacaaggcc aactgactg tagacacatc ctccagcaca 240
 10 gtttacatac aactcagcag cctgacatct gaggactccg cgggtctatta ctgtgcaaga 300
 ggggactatg gttgcccttt tgtttactgg ggccaaggca ccacggtcac cgtctccagt 360
 15 ggcggcgggcg gcagcgggtgg tgggtggttct gggggcgggcg gcagcgacat cgagctcact 420
 cagtctccat tctccctgcc tgtcagtctt ggaggtccag cctccatctc ttgcagatct 480
 agtcagagtc ttgtacacag taatagagac acttatttac attggttcct gcagaagcca 540
 20 ggccagtctc cagagctcct gatctacaga gtttccaacc gattttcttg ggtcccagac 600
 aggttcagtg gcagtggatc agggacagat ttcacactca agatcagcag agtggaggct 660
 25 gaggatctgg gagtttattt ctgttctcaa agtacacatg ttccattcac gttcggctcg 720
 gggaccaagc tggaaatcaa acgggcgggc gca 753

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<210> 17
 <211> 786
 <212> DNA
 <213> Artificial Sequence

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<220>
 <223> Description of Artificial Sequence: synthetic
 sequence substituting bacterial codons for mouse
 codons

40

<400> 17
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 tcttgccgcg cttctggcta caaattcacc acctactgga tgcaactgggt gcgccagcgc 120

45

-13-

cctggccaag gccctgagtg gattggcgat atttatcctg gtagtggtga ttctaactac 180
gatgtgaagt tcaagaacaa ggccacactg actgtagaca catcctccag cacagtttac 240
5 atccaactca gcagcctgac atctgaggac tccgcggtct attactgtgc aagagggggac 300
tatggttgcc cttttgttta ctggggccaa ggcaccacgg tcaccgtctc cagtggcggc 360
ggcggcagcg gtggtggtgg ttctgggggc ggcggcagcg acatcgagct cactcagtct 420
10 ccattctccc tgccgtgcag tcttggcgat ccagcctcca tctcttgccg ctctagtcag 480
agtcttgtag acagtaatcg cgacacctat ctgcattggt tcttgcagaa gccaggccag 540
15 tctccagagc tcttgatcta ccgcgtttcc aaccgctttt ctgggggtccc agaccgcttc 600
agtggcagtg gctcagggac agatttcaca ctcaagatca gcagcgtgga ggctgaggat 660
ctgggcgttt atttctgttc tcaaagtaca catgttccat tcacgttcgg ctcggggacc 720
20 aagctggaaa tcaaacgggc ggccgcaggt gcgccggtgc cgtatccgga tccgctggaa 780
ccgcgt 786

25

<210> 18
<211> 7
<212> PRT
30 <213> Homo sapiens

<400> 18
Gly Gln Asn Asp Thr Ser Gln
1 5

35